

Exhibit A

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10      20      30      40      50      60      70
SEQ ID NO:1 09/428,122  ---CGCTCTTACCTAGTAGAGGTTGA-GTGAATTTCTTGACTTGTTTCTCCTATTGGTGTATCTCTTAAAA
SEQ ID NO:1 09/376,770  -----
SEQ ID NO:3 09/376,770  -----
SEQ ID NO:5 09/376,770  -----
SEQ ID NO:7 09/376,770  -----
SEQ ID NO:9 09/376,770  -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 -----AAATGCTTCTAGTCTAGGGGAACCTTTTTC--TATTTCTACGAAGGGAGTGTCTT----G
SEQ ID NO:15 09/376,770 TTTGAAAAAAGAGAGTTACGGCAGCTGTAAAGTTTTTAA--TATTGCTCCCTTTGTTCCATTTATGTAGC

80      90      100     110     120     130     140
.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
SEQ ID NO:1 09/428,122  TATTAATTCAAAATCAAAGTATATATTTTACAATGAAGTCTTCTTTCCCAAGTTTGTATTTTCTACAT
SEQ ID NO:1 09/376,770  -----
SEQ ID NO:3 09/376,770  -----GGGAACCTTTTATTCGTTTTATTAAAGTAAGATCGAGTATGGTATTGAGAAACGTTCTCGT
SEQ ID NO:5 09/376,770  -----
SEQ ID NO:7 09/376,770  -----
SEQ ID NO:9 09/376,770  -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 TCTAGAACTCCATAGGGAGAT-CGCACGCTAAA-TATGAGAATATAATATGAAGACTTCAGTTTCTATGT
SEQ ID NO:15 09/376,770 GTTCAGACTTTGCACTAAAACGAGGGGTGTCATA-TGAGATCGTCTTTTCTTGTATTAAATATCTTCAT

150     160     170     180     190     200     210
.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
SEQ ID NO:1 09/428,122  TTGCTATTTTCCCTTTGTCTATGATTG--CTACCG-AGACAGTTTGGATTCAAGT--GCGAGTTTCGA
SEQ ID NO:1 09/376,770  -----
SEQ ID NO:3 09/376,770  AAAAGATGGTATTCTAGTAATTACAGTGAATTAACGACATATGTTAGGTTTCTTAAACGCTTCTTTGG
SEQ ID NO:5 09/376,770  -----
SEQ ID NO:7 09/376,770  -----
SEQ ID NO:9 09/376,770  -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 TGTGGCCCTGCTTTGCTCGGGGGCTAGCTCTATTGTACTCCATGCCGCAACCACTCCACTAAATCCTGA
SEQ ID NO:15 09/376,770 CTCTAGCCTTTCTCTCTTAATGAGTG--TTTCTGCAGATGCTGCCGATCTCACATTAGGGAGTCGTGA

220     230     240     250     260     270     280
.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
SEQ ID NO:1 09/428,122  TGGGAATAAAAAATGGTA-----ATTTTTCAGTTCGTGAGAGTCAG----GAAGATGCTGGAAC
SEQ ID NO:1 09/376,770  -----
SEQ ID NO:3 09/376,770  TTCTCTCAAGAGCGTATTCTAAAAAATTTCAAAAACCTGTAGATAAAGTGAACATTTATGATGAAATG
SEQ ID NO:5 09/376,770  -----
SEQ ID NO:7 09/376,770  -----
SEQ ID NO:9 09/376,770  -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 AGATGGGTTTATTGGGGAGGGCAATACAAATACTTTTCTCCGAAATCTACAACGGATGCTGCAGGAACT
SEQ ID NO:15 09/376,770 CAGTTATAATGGTGATACAAGCACCAAGAATTTACTCCTAAAGCGGCAACTTCTGATGCTAGTGGCAGC

290     300     310     320     330     340     350
.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
SEQ ID NO:1 09/428,122  ACCTACCTATTTAAGGGAATGTCACTCTAGAAAAATTTCTGGAACAGGCACAGCAATCACAAAAAGCT
SEQ ID NO:1 09/376,770  -----
SEQ ID NO:3 09/376,770  CTCACGCCTTTATCTGATGATGAATTGCGGAATAAGACTGCGGAATTAAAGCAGAGATATCAGAACGGAG
SEQ ID NO:5 09/376,770  -----
SEQ ID NO:7 09/376,770  -----
SEQ ID NO:9 09/376,770  -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 ACCTACTCTCTCACAGGAGAGGT--TCTGTATATAGATCCGGGGAAAGGTGGTTCAATTACAGGAACTT
SEQ ID NO:15 09/376,770 ACCTATATTCTCGATGGGGATGT--CTCGATAAGCCAAGCAGGGAA-ACAAACGAGCTTAACCACAAGTT

360     370     380     390     400     410     420
.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
SEQ ID NO:1 09/428,122  GTTTTAAACAACACTAAGGGCGATTGACTTTTACAGGTAACGGGAACCTCTATTGTTTCAACAGGTGGA
SEQ ID NO:1 09/376,770  -----
SEQ ID NO:3 09/376,770  AGTCTCTTGATAGCATGCTTCTGAAGCTTATGGTGTGCGTGAACAAACGTTTGTGCGACGTTTACAGGCAC
SEQ ID NO:5 09/376,770  -----
SEQ ID NO:7 09/376,770  -----
SEQ ID NO:9 09/376,770  -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 GCTTTGTAGAACTGCTGGCGATCTTACATTTTATAGGTAATGGAAATACCCTAAAGTTCTGTGCGGTAGA
SEQ ID NO:15 09/376,770 GTTTTCTAACACTGCAGGAAATCTTACCTTCTAGGGAACGGATTTTCTCTTCAATTTGACAATATTAT
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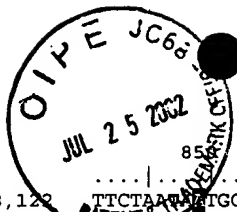
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8,122 01PE JC66-13
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TRADE MARK OFFICE
APR 28 1962 ACTGT

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850 860 870 880 890 900 910
 SEQ ID NO:1 09/428,122 TTCTAAGTGTCTAAGTTTCCTTTATTGACAA--TAAGGTCACAGG--AGCGAGCTCCT--C
 SEQ ID NO:1 09/376,770 ATGCTCAAAAGTTAAAGAAGGCCATGTGGTGGT--AGACTCTATCACAATCATAACGGA--AGGAGAAA
 SEQ ID NO:3 09/376,770 CATTGATGAAGTCGACTCGATCTTAATAGATGAAGCTAGAACACCCTTAATTATCTCAGGTCCTGGAGAA
 SEQ ID NO:5 09/376,770
 SEQ ID NO:7 09/376,770 TTATAATGA--ATAAGCTACTCAATTCGTCAG--CAGAACACTTGGTGGCGATACCGCCTTAAACATG
 SEQ ID NO:9 09/376,770
 SEQ ID NO:11 09/376,770
 SEQ ID NO:13 09/376,770 CACTGCGAACCATGAGTTGATATTTTCAAAAA--TAAGACTTCTGGGAATGCTGCAATGGCGGAGCC
 SEQ ID NO:15 09/376,770 CTCCAATAA--CCAAATATCTTTT--TCGAT-----GGCTGCAA-----AGCAA--C--T

920 930 940 950 960 970 980
 SEQ ID NO:1 09/428,122 AACAAC--GGGGGATATGTCAGGAGGTCTATCTGTGCTT-----ATAAACTAGTACAG-
 SEQ ID NO:1 09/376,770 ATGCTTCAAATAAACATCCCTTACCCAAATTAAGACCAGAAGTGGG--GCTCTTTTCTCAATTAGAT
 SEQ ID NO:3 09/376,770 AACATAATCCAGTCTATTTTGTAGCTTAAAGAAAAGTCGCAAGTCTAGTGATTTGCAAAAAGAGCTCT
 SEQ ID NO:5 09/376,770
 SEQ ID NO:7 09/376,770 ATCAATAAGTCCAGCGACTTAATCCTTGCTCTTTGGATGATGGGCGT--TGTCTTAATGATCAATTATTCC
 SEQ ID NO:9 09/376,770
 SEQ ID NO:11 09/376,770
 SEQ ID NO:13 09/376,770 ATAAATGCTCAGGGGACCTTACATTACTGATAACACTTCTTTGTT--ACTTCAAGAAAATAGCACAAT
 SEQ ID NO:15 09/376,770 ACAAAT--GGCGGAGCTATT--GATTGTAACAAAGCA-----GGGCGCAACCCAG-

990 1000 1010 1020 1030 1040 1050
 SEQ ID NO:1 09/428,122 ATACTAAGGTCACCCTCACTG-----GAAATCAGATGT--TAC-----TCTTCAGCAAC--
 SEQ ID NO:1 09/376,770 TTGTATGAAGACTTGAGAATTCTAGCTA--AGAATACGACTCTGTGAGCCTAAAGTAGAATTTCTGAA
 SEQ ID NO:3 09/376,770 GCAGCCGTATCGCATTAGAAGCACGTCGCGGCTTAGATAGCTTTTAGATGTTGATATCTTCTCTAAAGA
 SEQ ID NO:5 09/376,770
 SEQ ID NO:7 09/376,770 TTTGCTCCGCTATCGTTGACTTGATG--ATCACCATCAACTTATCGATCTCTGTATTCTTATTGATGGT
 SEQ ID NO:9 09/376,770
 SEQ ID NO:11 09/376,770
 SEQ ID NO:13 09/376,770 GCAGGATGGTGGAGCTTTGTGTAGCACA--GGAACCATAAGCATTACCGGTAGTGATTCTATCAATGTGAT
 SEQ ID NO:15 09/376,770 ACCCTATCTTGACTCTTTCAG-----GAAATGAGAGCC--TGC-----ATTTCTGAAT----

1060 1070 1080 1090 1100 1110 1120
 SEQ ID NO:1 09/428,122 ----AATACATCGACAACAGCGGGAGGAGCTATCTATGTGAAAAAGCTCGAACTGGCTTCCGGAGGACTT
 SEQ ID NO:1 09/376,770 GGGAAAACTAACATAGCCCTTACCTAATAGCTAAACCCTCAATTGAAATATTCATATCTCAGGAAATC
 SEQ ID NO:3 09/376,770 TAAAAAGTTCTTGAAGGCATCTCTGAATTTGCGCGAGCCTTTGGTTGGTAAGCAAGGAATGCCTTTG
 SEQ ID NO:5 09/376,770
 SEQ ID NO:7 09/376,770 GGCTCTTTATATTCCAAGTGCTTTGCGAGCTGTCTGTTTTCCCTCGTTGCTCTCATCACTACGATGTTT
 SEQ ID NO:9 09/376,770
 SEQ ID NO:11 09/376,770
 SEQ ID NO:13 09/376,770 AGGAAATACTTCAGGACAAAAAGGAGGAGCGATTCTGCGAGCTTCTCTCAAGATTTTGGGAGGGCAGGGA
 SEQ ID NO:15 09/376,770 ----AACACAGCAGGAAATAGTGGAGGTGCGATTATACCAAAAAATTGGTGTATCTCTCAGGACGAGGA

1130 1140 1150 1160 1170 1180 1190
 SEQ ID NO:1 09/428,122 ACCCTATTCAGTAGAAATAGTGTCAATGGAGGTACAGTCTCTAAAGGTGGAGCCATAGCTATCGAAGATA
 SEQ ID NO:1 09/376,770 AAGTCGTTCTGAACATAAAATTTTAAACCCCTACAAATTTACCGTAATGATCTCTTTGAACGAGAAAA
 SEQ ID NO:3 09/376,770 AATCGTGTGTACGTGCTGTACGTGAGCAGCCAGATCTTCGTGCTATGATCGATAAATGGGATGTTTATT
 SEQ ID NO:5 09/376,770
 SEQ ID NO:7 09/376,770 CGCTTGGGATTAATATTTCTCTCTCGACAGATTCTCTCTAAAGCGTATGCGAGTCATGTCATTACAGG
 SEQ ID NO:9 09/376,770
 SEQ ID NO:11 09/376,770
 SEQ ID NO:13 09/376,770 GGCGCTCTCTTTCTAATAACGTAGTGACTCATGCCACCCCTCTAGGAGGTGCCATTTTATCAACACAG
 SEQ ID NO:15 09/376,770 GGAGTGTTATTTCTAACAACAAAGCTGCGAATGCTACTCTCTAAAGGAGGGGCAATTGCGATTCTAGATT

1200 1210 1220 1230 1240 1250 1260
 SEQ ID NO:1 09/428,122 GTGGGAATTGAGTTTATCCGCCGATAGTGGTGACATTGTCTTTTAGGGAATACAGTCACCTCTACT--
 SEQ ID NO:1 09/376,770 ATTTCTTAAGGGTCTTGATGATCTAAGAACGTATTATCTCAAGCGAGGATATTTCCGATCCAGTGTAGAC
 SEQ ID NO:3 09/376,770 ATCATGCTGAGCAGAATAAAGAAGAGAGCCTAGAGCGTCTTTCAGAGCTCTACATTATTGTTGATGAGCA
 SEQ ID NO:5 09/376,770
 SEQ ID NO:7 09/376,770 CCTTCGGAGACTTCGTGGTTGGAGGGAACATATGTTGGTGGGTTCAATTCTTCTCTATTATTACAATCAT
 SEQ ID NO:9 09/376,770
 SEQ ID NO:11 09/376,770
 SEQ ID NO:13 09/376,770 GAGGATCCTTGCAGCTCTTCACTCAAGGAGGGGATATCGTATTCGAGGGGAATCAGGTCACTACAACAGC
 SEQ ID NO:15 09/376,770 CTGGAGAGATTAGATTTCTGCAGATCTCGGCAATATCATTTTCGAGGGCAATACTACGAGCACTACAGG

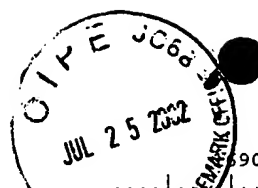
PIPE JC66 JUL 25 2002

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SEQ ID NO:1 09/376,770 -TACAGTCTGGAACACAATCAAGAAAAAGGTCACATCGATGTTTTAATTAAATCAATGAAGGTCCTTGC
SEQ ID NO:3 09/376,770 CAATAAGATTTTGTAGCTTACAGATAAAGGAATGCAGCAGTGGGTTGAGTATGCTGGAGGCTCTACCGAA
SEQ ID NO:5 09/376,770
SEQ ID NO:7 09/376,770 TCAGTTTATCGTAGTAATAAGGGTGCCGAGCGTGTGCGGAAGTTGCTGCCGATTCCGATTGGATGCG
SEQ ID NO:9 09/376,770
SEQ ID NO:11 09/376,770
SEQ ID NO:13 09/376,770 TCCAAATGCTACCCTAAGAGAAATGTAATTCACCTCGAGAGCACCGCGAAGTGGACGGGACTTGCTGCA
SEQ ID NO:15 09/376,770 AAGTCCGCGAGTGTGACCAGAAATGCTATAGATCTTGATCGAATGCAAAATTTTAAATCTCCGAGCG
1340 1350 1360 1370 1380 1390 1400
SEQ ID NO:1 09/428,122 GCTGCTGGTAGAGCCATCTACTTCTATGATCCCATAACTACAGGATCATCCACAACAGTTACAGATGTTCT
SEQ ID NO:1 09/376,770 GGGAAATTAACAGCTTACGTTCTCAGGAATCTCTCGATCAGAAAAATCAGATATCCAGAATTTATTTC
SEQ ID NO:3 09/376,770 GAGTTCGTGATGATGGATATGGGGCATCAGTATGCTCTTATAGAAAATGATGAGACCCATACCTTCGAG
SEQ ID NO:5 09/376,770
SEQ ID NO:7 09/376,770 ATGCCAGGTAAACAGATGGCGATTGATCGGACTTACGAGCTGGTATGATTGATGCCACACAAGTTCGTG
SEQ ID NO:9 09/376,770
SEQ ID NO:11 09/376,770
SEQ ID NO:13 09/376,770 AGTCAAGGTACGCTATCTATTCTATGATCCATTACCACCAAC--GAT-ACGGGAGCAAGCGATAACT
SEQ ID NO:15 09/376,770 ACTCGGGGAATAAAGTTATTTCTATGATCCTAT--C-ACGAGCTCAGGAGCTACTGATA-AGCTCTCT
1410 1420 1430 1440 1450 1460 1470
SEQ ID NO:1 09/428,122 TAAAAGTTAATGAGACTCCGGCAGATTCTGC---ACTACAATATACAGCGAAATCA-TCTTCACAGGA
SEQ ID NO:1 09/376,770 AAACCAAGCAGCACTCTACAACACAAGTTGGTTTACTGGAGCTGGACTCTATCAACCCAGATATTGTTGA
SEQ ID NO:3 09/376,770 ATAAGATCAATAAAAAAATGCAATTTCTGAAG--AAGACACCTTAAGAAAGGCTCGTCTCAGGATTA
SEQ ID NO:5 09/376,770 -----G-----AACAAAACGTTTGGCTTTCAC--TCTTGAGAAAA
SEQ ID NO:7 09/376,770 A-----TAAAGGGCTCAAATCCAAAAGGA---AAGTGAACTCTACGAGCCATGGACGGTCCCATGA
SEQ ID NO:9 09/376,770 -----GCTGTGATTTCTTAT-C---AAGAAATCTACCAGAAATCCGCATCCAGCCTTTA
SEQ ID NO:11 09/376,770
SEQ ID NO:13 09/376,770 TACGTATCAATGAGGTCAGTGCAAA-TC-----AAAAGCTCTCGGATCTATAG-TATTTCTGGA
SEQ ID NO:15 09/376,770 T-----TGAATAAAGCTGACGAGGATCTGG---AAATACCTATGAAGCTACATCG-TTTTCTCTGGA
1480 1490 1500 1510 1520 1530 1540
SEQ ID NO:1 09/428,122 GAAAAGTTATCAGAGACAGAGGCCCGCAGAT-TCTAAATACTTACTTCGAAGCTACTACAGCC---TGT
SEQ ID NO:1 09/376,770 ACAAGTAGCTTGCAATTACGAATACCTACATAATACGGGTACGCTGATGCTATAGTCAACTCTCAC
SEQ ID NO:3 09/376,770 CGACAGTTATTACGAGCCCACTTCTCATGGAGCTGTATGATGATTATATTGTCGCGACGATCAGATTG
SEQ ID NO:5 09/376,770 ACAAGGCATAAGGTAATTTGCAGATAA-T-T-CCGAATATAGTCTCTTAAATCGAACCTTGGC---AAC
SEQ ID NO:7 09/376,770 AGTTCTTCAAAGGAGACGTTATCGCTGGTA-T-CGTTTCTCTTTGATTAACATTGTTGGCGGTTT-GAC
SEQ ID NO:9 09/376,770 GGAAGGATTCAGATTTTCTATTGATACGT-TGTCGCTCATAGGAGGCATATGGCAGCATCAGGAGGCAC
SEQ ID NO:11 09/376,770
SEQ ID NO:13 09/376,770 GAGAGATTGTGACAGCAGAGCTATAG---CTGAATACTTACTTCGAGGATCAACCAGCC---TGT
SEQ ID NO:15 09/376,770 GAGAACTCTCAGAAGAGGCACTTAAGAAA-C-CTGACAATCTGAAGTCTACATTTACACAGGC---TGT
1550 1560 1570 1580 1590 1600 1610
SEQ ID NO:1 09/428,122 AACTCTTTCAGGAGTACTCTATCTTTAAACATGGAGTGACTCTG---CAGAGTCAGGCACTCACT
SEQ ID NO:1 09/376,770 TATGACCTTGACGACAAAGGAATATTCTTCTTACATGGATATTGATCGAGGGTCGCGATATACCTTAG
SEQ ID NO:3 09/376,770 TGATTATCGATGAACATACAGGACGTCCTCAACCTGGAGGAGCTTTTCTGAAGGCCTCCATCAAGCTAT
SEQ ID NO:5 09/376,770 AGCATGGGTGAAAA--AACAGAAAGGCCACGCCGAAGCCAGCTTAGA---GATGTCGGGAAAAAGGT
SEQ ID NO:7 09/376,770 GATTGGGGTGCTATGCACGGCATGGACTCGCTCAAGCAGTACAGTCTACACTCTTCTCTCATTTGGA
SEQ ID NO:9 09/376,770 AGGTGGTTTAGGAGGCACTCAGGGTGTCAACCTTGCAGCTGTAGAAGTGCAGCTGCAAAAGCAGATGCA
SEQ ID NO:11 09/376,770 -----GTAATCGTTTAGAT
SEQ ID NO:13 09/376,770 CACTTTAGTAGAGGGGAGCTTAGTACTTAAACAGGGAGTGACCTTG-----ATCACACAAGGATTCTCG
SEQ ID NO:15 09/376,770 AGAGCTTGCTGCAGGTGCCTTAGTATTGAAAGATGGAGTGACTGTA-----GTTGCAATATATTAACG
1620 1630 1640 1650 1660 1670 1680
SEQ ID NO:1 09/428,122 CACAGGCAGATTTCGCTCTCGAAATGGACGTAGGA-CTACTCTAGAACC---TGATGACTAGCACC
SEQ ID NO:1 09/376,770 GACACGTCCATATCAAGGGTTTGAGGTTTGCCAA--ACGCCTTATAGAAAAGCAATCCCAAGTCGGCC
SEQ ID NO:3 09/376,770 CGAAGCTAAAGAACAGCTCACTACCGTAAGGAATCTCAGACGCTGTGTACAGTCAAGTTGCAAAATTTTC
SEQ ID NO:5 09/376,770 CAACTAGCAAAATCTCAGGATTTTCTTCTGCGGTT-ACCTTTATCGTCTCTA-TGTTTACGGCTTTCTC
SEQ ID NO:7 09/376,770 GATGTTTAGTCTCTCAAAATCTCTCTTTTGATT-GCGTTGACAGCGGGTATTGTACGACTCGTGTA
SEQ ID NO:9 09/376,770 GCAGAAAGTTGTAGCAGCCAAGAAGGTTCTGAGATG-AACATGATCAACA-ATCTAGGACCTGACAAA
SEQ ID NO:11 09/376,770 GCTGTAAATATAACAATGAAGATATCCCAAAGC--ATCAGACTTCCCTAA-CCATATCTTGGTCAT
SEQ ID NO:13 09/376,770 CAGAGGCCAGAAATCAGCTTCTTTTGGATCTGGGG-ACCTCATTAAGGCTTCTAGAGAAGATATTGTC
SEQ ID NO:15 09/376,770 CAGTCTGAGGGATGAAAGTCGTATATGGATGGAGG-CTACTTTTGAGGCAAGCGGTGAGGGGCTCACT



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SEQ ID NO:1 09/428,122 ATAAATTTTGGTCTTAACATCAGTTCTATAGACGGTGCAAGAAGGCAAAATAGAAAACAAAGCTA
SEQ ID NO:1 09/376,770 TGAATGATCTTTATTGCCCCGATAA---AATATGGGATGGGGCTCATAAGATCAACAAACTTATGCAA
SEQ ID NO:3 09/376,770 TTCCGTCTATATGAAGAGCTTGCAGGGATGACCGGAACAGCAATACGGAGTCTCGAGAGTTTAAAGAAA
SEQ ID NO:5 09/376,770 CCTATCGACCTTTTTTTCAAGCA---TTTAGGTGGCTT---TCTGGTTTCCTTGCTCTCACAAGCTC
SEQ ID NO:7 09/376,770 TCGAGTGACAAAATACGAACCTGGG---TAAAGAGATTTCTACTCAGCTCGTTAAAGAACCACGAGCAC
SEQ ID NO:9 09/376,770 TCCCGCAGCAGCAACACGACGAAAAA---AAAAGGAAGAGAAGTTCAAACCTCTAGAATCTCGGAAAAAA
SEQ ID NO:11 09/376,770 GATTAAAAAAGGATTGC-CATGCAAAACCAATACGACCAATTACTAGAATCCTTAGCACCCCTATTAAAT
SEQ ID NO:13 09/376,770 ATCAACAAATTTATCTTAAATGCCGATACCATTTACGGAAGAAATCCTATCAATTTGTAGCTTCAGCAG
SEQ ID NO:15 09/376,770 CTCAATGGCCTAGCCATTATATATGATTCTTTAGATGGGACAAATAAGCTATCATTAAGGCGACGGCAG

SEQ ID NO:1 09/428,122 CGTCAAAAACTGACTTTTCTGGA-----ACCATC-ACTTTATTGGACCCGACG-GGCACGTTTAT
SEQ ID NO:1 09/376,770 AGTATGGCTACATCAATACCAATGTAGACGTTCTCTTCATCCCTACGCAACCCGCCCTATTATGATGT
SEQ ID NO:3 09/376,770 TTTATATCTTTATGTCCTCCAAGTACCCACGTTCAAGCCCTTGTACGCATAGATCATAATGATGAAT
SEQ ID NO:5 09/376,770 CCACTCGCCATGATCCTGTAATAC-----CTTATT-TTATTGAAGAACTGTCT-TATGCTTTCTT-T
SEQ ID NO:7 09/376,770 TACTCCTTGCAAGGTGCTGCAACTTTAGGG---GTTGGT-TTCTTCAAGGCTTCCCTCTATGGTCTTCT
SEQ ID NO:9 09/376,770 GGAGAACTGTTGAAAGGCTGAGAAAAAT---CTGAAT-CTACAGAAGAGAAGCTGACACAGATCTTGC
SEQ ID NO:11 09/376,770 CGACACTTGTCTCCAGATAAAAAATCACTCTGTTTAAT-CCGTTTCAGCGATACCCA-TGTCCCTGTG-C
SEQ ID NO:13 09/376,770 ACGAATAAGAACATTACCTTAACAGGA-----ACCTTA-GCACTGTAAATGCAGAT-GGAGCTTTCTAT
SEQ ID NO:15 09/376,770 CAAGTAAGGATGTTGCCTTATCAGGG-----CCTATC-ATGCTGTAGATGCTCAG-GGGAACATTAT

SEQ ID NO:1 09/428,122 GAAA---ATCATAG--TTTAAGAAATCCTCAGTCCACGACATCTTAGAGCTCAAA-----GCT
SEQ ID NO:1 09/376,770 AACTTATGAGTAAGTGAAGGGTCTCCTTATAAAGTGGGTTAATTAAATTAAGTGGGAATACCATACA
SEQ ID NO:3 09/376,770 TTAT---ATGACAGAGCGTGAGAAGTACCAGCTATTGTTAATGAGATCCGACTATTATGGAAGGGG
SEQ ID NO:5 09/376,770 TAAC---AGCATCA--CTTCCCTTACTGGGAGCTGTGTGCTGTTGTTGGCTCATGT-----A-GGT
SEQ ID NO:7 09/376,770 CCATTTTAGCATTAATTTTCTGCTGCTTAGGAGATTCTCCTACTGACTAAGAAATCAGCGG---CA-GGA
SEQ ID NO:9 09/376,770 TGATA--AGTATGCTTCTGGGAATTCTGAATCTCTGGTCAAGAACTTCCGGCTGCGT---GATGGA
SEQ ID NO:11 09/376,770 AAAT---AGAAGAAG-ATGGAATTCGGAGATCTTG--CAGTATCGACACTACTAG-----GTACT
SEQ ID NO:13 09/376,770 GAGA---ACCATAC--CTTGCAAGACTCTCAAGACTATAGCTTTGTAAAGTTATCTCCAG---GA-GCG
SEQ ID NO:15 09/376,770 GAGC---ATCATAA--TCTCAGTCAACAGCAGGTCCTTCTTTAATAGAGCTTTCT-----GCA

SEQ ID NO:1 09/428,122 TCTGGAAGCTTAACAAGCACCGCAGTGA-----CT-----CCAGATCCATAATCGG-----TGA
SEQ ID NO:1 09/376,770 AAATCTGACGTTATTTACAGGAACCAAGT---CTCTCCACAGAGATACATCAATCGCTTAAAGCTA
SEQ ID NO:3 09/376,770 AACCCATTCTTGTGGTACAGAATCTGTAGAGGTCTCTGAGAAGCTGTCTCGGATTTTACAGACAGAATC
SEQ ID NO:5 09/376,770 TTTCTTATCGTTGGTCTTACTTTTCTA-----CC-----GAAGTTTAAACAGCA-----TAT
SEQ ID NO:7 09/376,770 AAAAAGGTGGTGGCTCAGGAGCTTCAAA-----CAACCGTAGGGCTGCTGGTATGCGGCTGCTAC
SEQ ID NO:9 09/376,770 ATAGGAGACGATGCTTCTCCAGAAGACA-----TT-----CTTCTCTTTTACAAGAGAAAATTA-
SEQ ID NO:11 09/376,770 CTCTCTGAAAACGTTATTTCCGAGCGCTA-----TTTCAAACTCTCTCTCTGTAAA-----TGG
SEQ ID NO:13 09/376,770 GGAGGACTATAATTAATCTCAGATGCTT-----CT-----CAGAAGCCTTGAAGTAGCTCCTTC
SEQ ID NO:15 09/376,770 CAAGGAACGATGACTACTACAGATATCC-----CC-----GATACCCCAATTCTAAA-----TAC

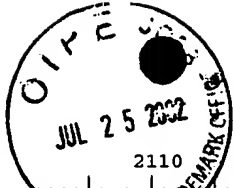
SEQ ID NO:1 09/428,122 GAAATTCATTACGGCTATCAGG-GAACTTGGGGCCC---AATTGTTTGGGG---GACAGGGGCTTCT
SEQ ID NO:1 09/376,770 GAAGATACTGAGCAAGCTTTAAG-AAATACAGGCTACTTCCAAAGCGTTAGTGTCTATACAGTTCTTCT
SEQ ID NO:3 09/376,770 GGATAGAGCATACTCTATTGAAT--GCTAAGAATCATGCTCAAGAAGCAGAAA-TTATAGCAGGAGCAGG
SEQ ID NO:5 09/376,770 CAAGAAGTTCAACC-CTATTGAG-AACATCAACA-----AACTTTAAAA-----TAAAGACTCTCAT
SEQ ID NO:7 09/376,770 TGTGGGGATAATCCGATGACT-ATTCTTAACCTCTCCGTAATTTAGAACTTGGAAAAAGATCTCTC
SEQ ID NO:9 09/376,770 -AAGACCCAGCTCTGCAATCCAC-AGCTTTGGACTAC---CTGTTCAAAC-----GACTCCACCTC
SEQ ID NO:11 09/376,770 CTCGTTCAATCCAGCATCAAGGAATTTAGGGTACG--GTGAGGCTACTCA---ACAGCTCTATCTTT
SEQ ID NO:13 09/376,770 TAGACCACATTTATGGCTATCAAG-GACATTGGAATGTGCAAGTCATCCAGG----AACGGGAACCTCAA
SEQ ID NO:15 09/376,770 TACGAATCACTATGGTATCAAG-GGAACGTGAAT-----ATTGTTTGGGT---CGACGATGCAACT

SEQ ID NO:1 09/428,122 ACCGAC-----TGCAACCTTCAA---CTGGACTAA-AACTGGCTATTTCTTAATCCCG
SEQ ID NO:1 09/376,770 CAACTTGATCCTATGGGCAATGGGATCAATACCGAGATATTTTGT-TAGAAGTCAAGAAACAACAACA
SEQ ID NO:3 09/376,770 GAAATTAG--GTGCTGTGACTGTAGCTACAAATATGGCTGGTGGAGGCAAGACATCAAACTAGATATG
SEQ ID NO:5 09/376,770 AGAGCT-----GATCAAAATCGATTTTAAAAATTTTGGAGCAG-CCTTAATTTTATACATAACGTTA
SEQ ID NO:7 09/376,770 TAAGCT-----TATCCCAACAGACAAATCAGGACAAAGCTT-TGTTGATGATGATGATCTCTAAAA
SEQ ID NO:9 09/376,770 CCAAG-----GTAAATTAAAA-----GAAGCGCT-TATCCAAGCAGGAATATCATA
SEQ ID NO:11 09/376,770 CAGAT-----ATCCTGAGTATGAA-----C--TACCTAAATGGAGAAAAATTTATT
SEQ ID NO:13 09/376,770 CCGAGC-----CAGGCAAAATTTAGA-----ATGGGTGCG-GACAGGATACCTTCCGAATCCCG
SEQ ID NO:15 09/376,770 GCATAAAA-----CAAAAATGCTACCTTAAC-----TTGGACTAA-AACAGGATACAGCCGAATCCAG

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2110 2120 2130 2140 2150 2160 2170

SEQ ID NO:1 09/428,122 ACCGCTCTCTCTCTTAGTCCCTAATAGCTTATGGAATGCATTATAGATA-TTAGCTCT-CTCCATT
SEQ ID NO:1 09/376,770 GGAAATTTAGGCTTATCTTAGGATTTAGTTCTCTTGACAATCTTTTGGAGGAATTGAAT-ATCTGAA
SEQ ID NO:3 09/376,770 AAGCTGTGATCGTAGGCGGTCTCTCATGTGATCGGTACTACACGGCATCAATCCCGTGGAAATTGATAGACA
SEQ ID NO:5 09/376,770 AAAAGC--AAAGTCTCTTTAATTATAGAACTGCAGGAGTCTCTCTATAAT-----TACT-GCTCAAA
SEQ ID NO:7 09/376,770 TGCGGG--AAGCTCTCTATCAGGATATCGGAATCCGATACCCTGGCAATCATG-TTCGACACA-GATTCCC
SEQ ID NO:9 09/376,770 CGGAGC--AATTCGGACGAACCTGCTATTGGTGCAGAAAACATCTTATTGCGCT-CTC-AAGA-ATATGCA
SEQ ID NO:11 09/376,770 CG-AGT--ATCTCAAGCTCTTTCTTTGCTATGCTAAGA--TTTGATGGAAT--CCCTAAGA-ACAGGGA
SEQ ID NO:13 09/376,770 AACGGC--AAGGATCTTTAGTTCCCAATAGCCTGTGGGTCTTTTGTTCATC-AGCGTGCT-ATCCAAG
SEQ ID NO:15 09/376,770 AACGTC--AGGGACCTTTGGTCTCTAATAGCCTGTGGGTCTTTGTTCGATG-TCCGCTCC-ATTTCAGA

2180 2190 2200 2210 2220 2230 2240

SEQ ID NO:1 09/428,122 ATCTTATGAG--ACTGCAACGAAGGGTTGCAGGGAGACCGTGCTTTTGTGTG-CTGGATTATCT
SEQ ID NO:1 09/376,770 AGTAATTTTATCTATTGAGAGTAGAAATATATTTCTAAAGGTTTTCGTTCTCTA-AGAGCGGTGGA
SEQ ID NO:3 09/376,770 GTTCCGCTGGCGGTCTGCTCGTTTAGGAGACCTGGCGCTGCGAAATCTTTTATCTTTTGAGATCGG
SEQ ID NO:5 09/376,770 TCTTCAAGAGAA--TTTTTATTAAGCAACCTCGATAGGAATTTTCTTTTATTGTTGGATTCTT
SEQ ID NO:7 09/376,770 CTCTTTTAAAG--GATACGATTATATGATCTGCTTAATGAAGTCCCTTATGTGCGAGAGAAATTCCT
SEQ ID NO:9 09/376,770 GACCAACTGAAT--GTTTCTCTTCAAGGGCTTCGCTCTTGTA--CTTGAAGTGA--CTGGAGACACA
SEQ ID NO:11 09/376,770 ATCTTCTTGACC--TTCATGTTTGGGAATCTACTACGTCGCG--TGAATGTT--TAAATACACA
SEQ ID NO:13 09/376,770 AACTCATGTAA--ATAGTCCCAAGCTTATGTCAGGAACGGGAGTCTGGGAG--CTCGAATTCG
SEQ ID NO:15 09/376,770 GCCTCATGACC--GGAGCACAAGTTCGTTATCTTCGTCAACAAATTTGTTGTTAT--CAGCAATCCG

2250 2260 2270 2280 2290 2300 2310

SEQ ID NO:1 09/428,122 A-ACCTCTTCCATAGGATAGTACAAAACACGA-----CGCGGGTTTC--GCCATTGAGTGGCGGT-
SEQ ID NO:1 09/376,770 GAACATCTATTCTTAAAGCCAACTTCGGGAGCA-----AAGTCAAGACTATACCTTTGAAGTGGACCAA
SEQ ID NO:3 09/376,770 CTGATCGGACTATTCGCCTCCCTCAAAATGAATACCCTTATCCGTCAATTTCTCTCCAGAGGAGAGG
SEQ ID NO:5 09/376,770 G-ACCTTGCTATCAGCGCC--ACAATTTCCGTA-----AAGAATTAA--GATGGAGAAGTTTGAG-
SEQ ID NO:7 09/376,770 CCGCAACCATGTGTTAAC-CA--ATGAGGTGG-AG-----GACATCTCA--CCGTTATATATCTACCT-
SEQ ID NO:9 09/376,770 C-ATACCTGTGATCAGCTACTTTCTATGCTTCA-----GACCGCTATA--CCTACCAAGATATGCT-
SEQ ID NO:11 09/376,770 AAACACTCACCCTCAGC-----AC-ATGCTTGA-----AACTTATAGGAACCTCTCTTAACACGGGA-
SEQ ID NO:13 09/376,770 A-ATTTCTACATAGAGAT--AAAATTAATGAG--CAGCGCTATC--GCCATAGCGGTGTCGCT-
SEQ ID NO:15 09/376,770 G-ACITTTTGCATCAGATCAGAAAGGAACCA-----CGTAGTTATC--CTCATCTAGCGCGGT-

2320 2330 2340 2350 2360 2370 2380

SEQ ID NO:1 09/428,122 -----T-ATGTATAGAGGAACCTACATACTT--GTTCAAGATAAGATTCTTAGTGCTCATTGTG
SEQ ID NO:1 09/376,770 ACCTCATTTCTAAACACTCCTTGGATTTTAGCAATTGAATTAGATAAATCAATTACAGACATTATCT
SEQ ID NO:3 09/376,770 CGATGTCGGACCCCTATGTTTAAATAGACTCATACAAACAGCACAGAAACGTGTGGAAGGGAGAACTATAC
SEQ ID NO:5 09/376,770 -----GTTAAGCAGGAGTTTAAAGACACCGA-----AGGAATCCTGAGATTAAAGCCCGTCGT
SEQ ID NO:7 09/376,770 -----TTTATTACCTATAAG--AATGCTGCGGT--CT--TCCTTCTGCTGGGTATAGTCAAGAT-GC
SEQ ID NO:9 09/376,770 -----ATTGTCAGCTCCTTCT--AATGAAAGCAAT--GGCAACAGATTAAAAGGCAGGCTCCCT-AC
SEQ ID NO:11 09/376,770 -----TTTATCTCCACTATTTTCAATACACAAA--A-----
SEQ ID NO:13 09/376,770 -----T--ATCTGTGGGAGTTGGCACTCATCCTT--TTTCTGATGCTACGATAAATGCGCTTTTGTG
SEQ ID NO:15 09/376,770 -----T-ATGCATTAGAGGAGGATTCTTACGG--CTTCTGAATTTCTTTAATTTTCTTTTGT

2390 2400 2410 2420 2430 2440 2450

SEQ ID NO:1 09/428,122 CAGCTCTTTGGAAGAGATAGAGACTACT-TTGTAGCTAAGATCAAG-GTACAGTCT--ACGGAGGAAC-
SEQ ID NO:1 09/376,770 AAAGATTATGCTGTCCAAACCTATGGCGGGAACGTGACACACAGTATATCTTGAACGAACAGCTGAAAT
SEQ ID NO:3 09/376,770 TATCCGTAAAGCATACCTTAGAGTATGATGATCATGAATAACGAGAGACAAGCGATATACCGTTTCCGC
SEQ ID NO:5 09/376,770 CGACAAATGCTCAAGAAATTCCCTA--TGAAGACTCGTCTACACA-G-GTGAAC--ATGCAAGCAC-
SEQ ID NO:7 09/376,770 AAAAGCTATTCTAGAGAGGCGACGAAT--TAAATATTGGACGCGCTCGAAGTGATC--ATTCTCCATCT
SEQ ID NO:9 09/376,770 GTACCCAGTGGCAACTACAAGTTC--TCATGACAGAACTCGTAACCTGCAAGC--AGTCTTACC--
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 CAGCTCTTCAGTAGAGATAAAGACTACG-TAGTATCAAAAATCATG-GAAGTAGCT--ACTCAGGGGTC
SEQ ID NO:15 09/376,770 CAGCTTTTGGCTACGACAGSACCATC-TTGTGGCTAAGAACCATATA-CCCATGTAT--ATCAGGGGC-

2460 2470 2480 2490 2500 2510 2520

SEQ ID NO:1 09/428,122 -TCTCTATTACCAGCACAAACCACTATATCTCTTCC--TTGCAAACTACGGCCTTGTTCGTTGTCT
SEQ ID NO:1 09/376,770 ACGGTCTATTTTATCGAGGAAGTCAAACGAGTTTACATGAAAAACGTAAGTTCTCTCCTAGGCCAATAT
SEQ ID NO:3 09/376,770 CATGATGCTTACATGCGGAATCTGTTTTCGATCTTGCAAAAGAAATCTATGCCATGTGTCTCTGATGG
SEQ ID NO:5 09/376,770 --CGTAGTCTCTAATCCCAAAGATATGCTGTGTCTAT--TG--GCTACATGCCTG--AAAATAT
SEQ ID NO:7 09/376,770 TTCCGATTTTCCATAAAAGCTCTCAAGATTTTGGGAATTCAAGAGGTACGTTCTATGATCGAATTT
SEQ ID NO:9 09/376,770 -TCGTACGATTACTTTGAAAGTCGCGTTCCTATTTTAC--TCGATAGCTTAAAAGCTG--AGGGAATC
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 GTATTTCTGAGGATACCTAGAGTTTGAAGTCCACAGGGATTCTATCTGATAGCTCCTCAGAAGCTT
SEQ ID NO:15 09/376,770 -AATGAGTTACCGACACCTCGGAGAGTTAAGACCCTC-----GCTAAGATTTGTCAGGAATTT

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2530 2540 2550 2560 2570 2580 2590

SEQ ID NO:1 09/428,122 ATATCTCTTACA-GAGA--TTCCTGTCTCTTTTCAGGAA----ACCTAGCTACACCCATAC----G
SEQ ID NO:1 09/376,770 AGACAGCAATAAAGGAT--TTGTCTCTGCTGCAGGTGTCAACTTGAATTACGATTCTGTAGATAGTCCTA
SEQ ID NO:3 09/376,770 TAGCATCTCTTAGTGATGAGTGATCTGTCAGTTCAAAGGGTGGACATTGCCAAATCTTGAAGAATGGATAAC
SEQ ID NO:5 09/376,770 AAAGCACCTTGG--GATCATTTGCCATGGGCATCA----ACTTACG--AGCTAAAA--G
SEQ ID NO:7 09/376,770 ATGGAACTTTCA-TTCC--CGGACTTAGTGAAGGAAGTCACA-AGGCTTATTCCATTGCAAAAGCTTACG
SEQ ID NO:9 09/376,770 CAAACTCCTTCT-GATC--TAAACTTTGTGAAGGTAGCTG--AGTCCATACCATAAAATCATTAAACGATA
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 GCTGTAAACCA-----AGTCTGCTACTATAGATATGC-----AGTTGCTTACAGCCATAG-----A
SEQ ID NO:15 09/376,770 CTGACTCCC-----TACCTTTTGTCTTCAATGCTC-----GGTTTGTCTTATGGCCATAC-----C

2600 2610 2620 2630 2640 2650 2660

SEQ ID NO:1 09/428,122 GATAACGAT---CTGAAAACCAAGTA-TACAACATATCCT-ACTCTTAAAG--GAAGCT---GGGGGAAT
SEQ ID NO:1 09/376,770 GAACTCCAACACTACAGGATTGCGGGGGGGGTGACTTTTGAGGTTTCTGGTT--TGGGAG--GAACTTAT
SEQ ID NO:3 09/376,770 CTCATCTTTCCCAATAGCCTTAAATATAGAAGAACTCAGACAGCTTAAAGATACAGATTCTATTGTCTGAA
SEQ ID NO:5 09/376,770 GATACTTG---ATGAAGCTGAAAG-TACGGAATTCOC---ATTATGC--GAAAC---GTACCTTT
SEQ ID NO:7 09/376,770 GAAATCTTT---AAGAGATTGGTTCA-AGAGCAATCTCA---ATTA---AAGAC---CTACGTAC
SEQ ID NO:9 09/376,770 AGTTCCCAA---CAGCATCTAAAGTAGAACGAGAAGTCCGCAATCTCATAG--GAGACG--ATGTTGAT
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 AATAATGAT---ATGAAAACCAAAAT-CACGACATATCA-GAAGCTCAGG--GATCTT---GGGCAAT
SEQ ID NO:15 09/376,770 GACAATAAC---ATGACCACAAAGTA-CACTGGCTATTCT-CCTGTTAAGG--GAAGCT---GGGGAAAT

2670 2680 2690 2700 2710 2720 2730

SEQ ID NO:1 09/428,122 GATAGTTTCCCT--TTAGAATTCGGTGAAGAGCTCCGAT--TTGCTTAGATGAAAGTGCTCT-----A
SEQ ID NO:1 09/376,770 CATTTTACAAA--CTCTCTTTAAACAGCTCTATCTATAGAA-AACTTACGCGTAAAGGTATTT-----T
SEQ ID NO:3 09/376,770 AAGATCGCTCTGAATTGATTCAGAGTTTCAAGTACGCTTTGATCATATGGTAGAAGGGCTTCCCAAAG
SEQ ID NO:5 09/376,770 AGCAGATCAGCT--TTTGGATGAAGGCAAGGAATAAAT--TTATTCCAG-AA-----TCT-----A
SEQ ID NO:7 09/376,770 AATCTTAGAATC--TCTAGCGAGTGGGCCCAACTGAGA--AAGATACAG-----TTTT-----G
SEQ ID NO:9 09/376,770 TCTGTGACCCCTGCTTGACTTATTCTTTTCTGCTTTACGT-CAAACGTCGTACGCCTTTTC-----T
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 GATGTTTTTCT--CTTGACTTTGGAGCGACTACAT--ACTA-CTACCCTAACAGTACTTTTT-----A
SEQ ID NO:15 09/376,770 GATGCCTTCCCT--ATAGAATGTGGAGGAGCTATCCCGGTAG-TTGCTTCAGGACGTCGGTCTT-----G

2740 2750 2760 2770 2780 2790 2800

SEQ ID NO:1 09/428,122 TTTGAGCAGT-TCATGCCCTTCATG--AAATTGCAGTTTG--TCTATGCACATCAGGAAGGTTTTTAA
SEQ ID NO:1 09/376,770 GAAAATCAAAGGGGAAGCTCAATTTATTAACCTATAGCAA-TACTACAGCTGAAGGAGATCCTGTGAG
SEQ ID NO:3 09/376,770 CTGGAGGGGAAGAAATGGATGCACTGCTATTGTAGAGATGTCGTTCCGTTCTGTATGGTATGCAAT
SEQ ID NO:5 09/376,770 CTTACGAAGCTATTGGAGAAATCT--ACTCTATATCACT--TCACTGAATGCGCAAAATCCTAATA
SEQ ID NO:7 09/376,770 CTTACAGAAT-ATGTACGGTCTTCTTTAAGAGCTTATATCA--GCTTCAAGTTCTCTCAAGGACAATCA
SEQ ID NO:9 09/376,770 CTTACAGACAGCAACGTCAGCAATTAGGAGCTATGATTGCTAATGCTTATAGATGCTGTAAATATAAACA
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 TTTGATTACT-ACTCTCCGTTTCTC--AGGCTGCAGTGCA--CCTATGCTCACCAGGAAGACTTCAAA
SEQ ID NO:15 09/376,770 GGTGGATACCCACAGCCATTCTA--AACCTAGAGATGA--TCTATGCACATCAGAATGACTTTTAG

2810 2820 2830 2840 2850 2860 2870

SEQ ID NO:1 09/428,122 G-AACAGGGAACAGAGCTCGTGAATTTGGAA-GTAGCCGCTCTGTGAATCTTCC--TTACCTATCGG
SEQ ID NO:1 09/376,770 TGAGCGCTTCTCTCTAGGTGGAGAGACTACAGTTCCGGGATATAAATCCTTTATTATCGGTCCAAAATAC
SEQ ID NO:3 09/376,770 T-GATGAGCAGTGGCGGATTCATCTTTAGATATGGACTTACTACGGAGTGAAGTTGGCCTACGTACTGT
SEQ ID NO:5 09/376,770 T-AAAAACTAACAACCTGATCATTTATAA-TGAATAAGCTACTCAATTTCC-----TCAGCGAAC
SEQ ID NO:7 09/376,770 --GCAATTTCTGTTTATCTCTTAGATCCAGAAATTGAAGAGATGATTCTGTGGAGCA--ATTAAACAGAC
SEQ ID NO:9 09/376,770 T-GAAGATTATCCCAAGCATCAGACTTCCCT--AAACCCTATCTTGGTCATG-----ATTAAAAAGG
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 G-AGACAGGAGGTGAGGTTCTGTCACCTTACTA-GCGGAGATCTTTTCAATTTAGCA--GTTCTTATTGG
SEQ ID NO:15 09/376,770 G-AAAACGGCACAGAGGCGGTTCTTCCAAA-GTGAAGACCTTCAATCTAGCG--GTTCTGTGAGG

2880 2890 2900 2910 2920 2930 2940

SEQ ID NO:1 09/428,122 GATCCGATTTGATAAGGAATCAGACTGCCAAGA-TGCAACGTAACAATCTAATCTTGGTTTACTGTGGA
SEQ ID NO:1 09/376,770 TCTGCTACAGAACCTCAGGAGGACTCTCTTCTGCTCTTATTTCAGAAGAGTTTCAATACCCTCTCATCA
SEQ ID NO:3 09/376,770 AGGGCAAAAAGATCCTTTGTAGAAATTTAAACA-CGAGTCTTTCTTACTGTTTGAAGAGCTTGATTCTGTA
SEQ ID NO:5 09/376,770 ACTTGGTGGCGATACCGCTTAAACATGATCAA-TAAGTCCAGCGACTTAAATCCTTGCTCTT-TGGATGA
SEQ ID NO:7 09/376,770 ATCGGCAAGTTCT--TACCTTGCTCTAGATCC--TGATTCTGTGAACCTAAATTTAAATCTATGAGGAA
SEQ ID NO:9 09/376,770 ATTGCCATGCAAAACCAATACGAGGAATTACTA-GAATCTTAGCAGCCC-TATTAATACGACACTTGC
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 CGTGAAGTTTGAGAGATTTTCACTGTAAAAG-GGGATCTTATGAACCTACCCCTTGCTTATGTTCTCTGA
SEQ ID NO:15 09/376,770 GATAAAATTTGA---GAAAT---CTCCGATAA-GTCTACGATGATCTCTCCATAGCTTACGTTCCCGA

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2960 2970 2980 2990 3000 3010

SEQ ID NO:1 09/428,122 PATENT 6,140,240
SEQ ID NO:1 09/376,770 TCGTAGTAACCCCGACTGTACGACAACACTGCGAATTAGCGGTGATTCTTGGAAAACCTTCGG
SEQ ID NO:3 09/376,770 GACAACCTAATAATTAGTGCCCTTGTATTCTTAGACTCAGGTTTGTGCGGTTTACAAGAGTATAAGATTTC
SEQ ID NO:5 09/376,770 TATTC-GTATTACGATTCCGCGGCATCTTTCCGCTCTGAACTGACCGTAGAGCCTAATCCTCGTGTCAA
SEQ ID NO:7 09/376,770 TGGGC-GTGTCTTAATGA-
SEQ ID NO:9 09/376,770 TACG--ATCACGCCAACACCTGCAGGAGGCCAACACCAGTATTATTACAGCAATTGATGTAAGAAGAT
SEQ ID NO:11 09/376,770 TCCAG-ATAAAATAACTCTTGT-----
SEQ ID NO:13 09/376,770 TGTGA-TTCGCAAGATCCCAAGAGCACGGCAACA-TT--GGCTAGTGGAGCTACGTGGAGCACCCACGG
SEQ ID NO:15 09/376,770 TGTGA-TTCGTAATGATCCAGGCTGCACGACAACCTTATGGTTTCTGGGATTCTTGGTCGACATGTGG

3020 3030 3040 3050 3060 3070 3080

SEQ ID NO:1 09/428,122 TACGAATTTGGCAAGACAAGCTTTAGTCCTTCGTGCAGGGAACCTTTTGTCTTAACTCAAATTTTGAA
SEQ ID NO:1 09/376,770 GTTAAAAGATCTACGTAGTAGTGTGGATTGGTCTGCGCTTCGATGTAATGAATAATGTTCTGTATTG
SEQ ID NO:3 09/376,770 CAACGTGATTCTACTGTAGCTACGCTTTTCATAATAATGTGAATTACGGTCCGTTAGAGTTGACTGTA
SEQ ID NO:5 09/376,770 ATGTACGAAAATTAATAGAAACAGAAATCCCTGACATTGCTGTGTTTCTTATCAAGAAATCCTACCAGA
SEQ ID NO:7 09/376,770 -----
SEQ ID NO:9 09/376,770 -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 AAACAATCTCTCCAGACAAGGATTACAACCTGCGTTTAGGGAACCTGTCTCATAAATCCTGGAATTGAG
SEQ ID NO:15 09/376,770 TACAAGCTTGTCTAGACAAGCTCTTCTGTACGTGCTGGAAATCATGACCTTTCCTTCAAACCTTTGAA

3090 3100 3110 3120 3130 3140 3150

SEQ ID NO:1 09/428,122 GCCTTTAGCCAAATTTCTTTTGAATTGCGTGGGTTCATCTCGCAATTACAATGTAGACTTAGGAGCAAAAT
SEQ ID NO:1 09/376,770 TTAGGATTTGGTTGGCCCTTCCGTCCAACCGAGACTTTGAATGGAGAAAAATGTATCTCAGCGAT
SEQ ID NO:3 09/376,770 GTTACAGATTCTGAAGATCAAGATTAAGAAATAACCTAGAAGGAGGAGTCTTCTAGGTAATTGTTTCTC
SEQ ID NO:5 09/376,770 AATCCGCATCCAGCCTTTAGGAAGAATTGAGATTCTTAATTGATACGTTGTCGCTCATAGGAGGCATGT
SEQ ID NO:7 09/376,770 -----
SEQ ID NO:9 09/376,770 -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 GTGTTTCAGTCACGGAGCTATTGAATTGCGGGGATCCTCTCGTAATTATAACATCAATTCGGGGGTAAAT
SEQ ID NO:15 09/376,770 GTTTTCAGTCAGTTTGAAGTCGAGTTGCGAGGTTCTTCTCGTAGCTATGCTATCGATCTGGAGGAAGCT

3160 3170 3180 3190 3200 3210 3220

SEQ ID NO:1 09/428,122 ACCAATTCTAATGCGTTAGCTTTGGTAAAGAGCTCCATACATCGAAGGAAAAAGAGCT--TTT--AAG
SEQ ID NO:1 09/376,770 TCTTCTTTGCTTTAGGGGGCATGTCTAAGATATAAATTAAGGACTTATCGAAGGAAATCTTTGTTGTTT
SEQ ID NO:3 09/376,770 AGGCTACATAGTTTTTATTTTAAATAAATGGGGTCCATCTGACCAACTTGGTCATT--ATAGTAGCG
SEQ ID NO:5 09/376,770 GGCAGCATCAGGAGGCACAGGTGGTTTAGGAGGCACTCAGGGTGTCAACCTTGAGCTGTAGAAGCTGCA
SEQ ID NO:7 09/376,770 -----
SEQ ID NO:9 09/376,770 -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 ACCGATTTTAATAGGGAAGTGAAGTTCCTTATTGTAGAGAGTATTTACAAGGATTTTGAGACGAAAGAG
SEQ ID NO:15 09/376,770 TCGGATTTTAATCC--TAAGTTTTCAACGAGATAGCATCAGGGTAAGCCAGGGCTCT--ATGTAAGAG

3230 3240 3250 3260 3270 3280 3290

SEQ ID NO:1 09/428,122 ATTTCTTGA--AGGCTCTTTTCGATT--TCGATTTCCATTTTAGTGTTTTGCTAAACACTTTC--
SEQ ID NO:1 09/376,770 TCAGAAAAGGCTTTTGGTACCCTTTTCTTATACCCAAAGTTAGT--
SEQ ID NO:3 09/376,770 AATCCAGAATTTCTGCGATTGCCGCTATGAGGAAGTGGCATAGTGCTCAGTGTATTGTTTGCTATA
SEQ ID NO:5 09/376,770 -----
SEQ ID NO:7 09/376,770 GCTGCAAAAGCAGATGC-----
SEQ ID NO:9 09/376,770 -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 ATTTCTTGTGTAGGTATCTCTCCACAGGGTCTAGTTCAATTTCTATAGTTCA--
SEQ ID NO:15 09/376,770 ATTTTCATAG--AGCCCTCTCTTTGTC--TTGCTTTTGTATTTTATTTT--

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SEQ ID NO:1 09/428,122 -----
SEQ ID NO:1 09/376,770 -----
SEQ ID NO:3 09/376,770 CAGAAA
SEQ ID NO:5 09/376,770 -----
SEQ ID NO:7 09/376,770 -----
SEQ ID NO:9 09/376,770 -----
SEQ ID NO:11 09/376,770 -----
SEQ ID NO:13 09/376,770 -----
SEQ ID NO:15 09/376,770 -----